RAW SEQUENCE LISTING ERROR REPORT
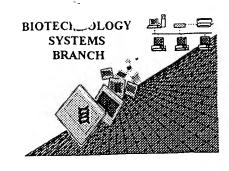
The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: *09/723,326*

Source: *O/PE*

Date Processed by STIC: *5-23-01*

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).
Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

<u>Checker Version 3.0 can be down loaded from the USPTO website at the following address:</u>
http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _09/723, 326_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1_____Wrapped Nucleics Wrapped Aminos — The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2_____Invalid Line Length — The rules require that a line not exceed 72 characters in length. This includes white spaces.

3_____Misaligned Amino Numbering — The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4_____Non-ASCII — The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5_____Variable Length — Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6_____PatentIn 2.0 "bug" — A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7_____Skipped Sequences (OLD RULES) — Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8_____Skipped Sequences (NEW RULES) — Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9_____Use of n's or Xaa's (NEW RULES) — Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10_____Invalid <213> Response — Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 ✓ Use of <220> — Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12_____PatentIn 2.0 "bug" — Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

AMC – Biotechnology Systems Branch – 06/04/2001

RAW SEQUENCE LISTING   DATE: 05/23/2001
PATENT APPLICATION: US/09/723,326   TIME: 10:32:47

Input Set : A:\09723326.txt
Output Set: C:\CRF3\05232001\I723326.raw

```
   3 <110> APPLICANT: WEBSTER, Keith A.
   5 <120> TITLE OF INVENTION: A Molecular Switch for Regulating Mammalian Gene Expression
   7 <130> FILE REFERENCE: 70373/268461 - UM99-29
   9 <140> CURRENT APPLICATION NUMBER: US 09/723,326
  10 <141> CURRENT FILING DATE: 2000-11-28
  12 <150> PRIOR APPLICATION NUMBER: US 60/171,597
  13 <151> PRIOR FILING DATE: 1999-12-23
  15 <160> NUMBER OF SEQ ID NOS: 16
  17 <170> SOFTWARE: PatentIn version 3.0
  19 <210> SEQ ID NO: 1
  20 <211> LENGTH: 21
  21 <212> TYPE: DNA
C--> 22 <213> ORGANISM: Artificial
  24 <220> FEATURE:
  25 <223> OTHER INFORMATION: oligonucleotide
  27 <400> SEQUENCE: 1
  28 tgtcacgtcc tgcacgacgt a
  31 <210> SEQ ID NO: 2
  32 <211> LENGTH: 22
  33 <212> TYPE: DNA
C--> 34 <213> ORGANISM: Artificial
  36 <220> FEATURE:
  37 <223> OTHER INFORMATION: oligonucleotide
  39 <400> SEQUENCE: 2
  40 cttcagcacc gcggacagtg cc
  43 <210> SEQ ID NO: 3
  44 <211> LENGTH: 22
  45 <212> TYPE: DNA
C--> 46 <213> ORGANISM: Artificial
  48 <220> FEATURE:
  49 <223> OTHER INFORMATION: oligonucleotide
  51 <400> SEQUENCE: 3
  52 tgtccattcc tgcacgacgt ac
  55 <210> SEQ ID NO: 4
  56 <211> LENGTH: 22
  57 <212> TYPE: DNA
C--> 58 <213> ORGANISM: Artificial
  60 <220> FEATURE:
  61 <223> OTHER INFORMATION: oligonucleotide
  63 <400> SEQUENCE: 4
  64 cttcagcacc gcttacagtg cc
  67 <210> SEQ ID NO: 5
  68 <211> LENGTH: 43
  69 <212> TYPE: DNA
C--> 70 <213> ORGANISM: Artificial
  72 <220> FEATURE:
  73 <223> OTHER INFORMATION: oligonucleotide
```

Does Not Comply
Corrected Diskette Needed
global error

Circled <223> response
must be more specific.
What is the source of
the genetic material in
the artificial sequences?
See # 11 on the Error
Summary Sheet.

Note: This error is indicated
throughout the sequence
listing. please review
and correct.

RAW SEQUENCE LISTING   DATE: 05/23/2001
PATENT APPLICATION: **US/09/723,326**   TIME: 10:32:47

Input Set : **A:\09723326.txt**
Output Set: **C:\CRF3\05232001\I723326.raw**

```
75 <400> SEQUENCE: 5
76 cttcagcacc gcggacagtg cctgtcacgt cctgcacgac gta                43
79 <210> SEQ ID NO: 6
80 <211> LENGTH: 86
81 <212> TYPE: DNA
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C--> 82 **<213> ORGANISM: Artificial**
```
84 <220> FEATURE:
85 <223> OTHER INFORMATION: oligonucleotide
87 <400> SEQUENCE: 6
88 cttcagcacc gcggacagtg cctgtcacgt cctgcacgac gtacttcagc accgcggaca   60
90 gtgcctgtca cgtcctgcac gacgta                86
93 <210> SEQ ID NO: 7
94 <211> LENGTH: 123
95 <212> TYPE: DNA
```
C--> 96 **<213> ORGANISM: Artificial**
```
98 <220> FEATURE:
99 <223> OTHER INFORMATION: oligonucleotide
101 <400> SEQUENCE: 7
102 cttcagcacc gcggacagtg cctgtcacgt cctgcacgac ttcagcaccg cggacagtgc   60
104 ctgtcacgtc ctgcacgact tcagcaccgc ggacagtgcc tgtcacgtcc tgcacgacgg  120
106 tac                123
109 <210> SEQ ID NO: 8
110 <211> LENGTH: 33
111 <212> TYPE: DNA
```
C--> 112 **<213> ORGANISM: Artificial**
```
114 <220> FEATURE:
115 <223> OTHER INFORMATION: oligonucleotide
117 <400> SEQUENCE: 8
118 cttcagcacc gcggacagtc acgtcctgca cga                33
121 <210> SEQ ID NO: 9
122 <211> LENGTH: 129
123 <212> TYPE: DNA
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C--> 124 **<213> ORGANISM: Artificial**
```
126 <220> FEATURE:
127 <223> OTHER INFORMATION: oligonucleotide
129 <400> SEQUENCE: 9
130 cttcagcacc gcttacagtg cctgtcacgt cctgcacgac gtacttcagc accgcttaca   60
132 gtgcctgtca cgtcctgcac gacgtacttc agcaccgctt acagtgcctg tcacgtcctg  120
134 cacgacgta                129
137 <210> SEQ ID NO: 10
138 <211> LENGTH: 132
139 <212> TYPE: DNA
```
C--> 140 **<213> ORGANISM: Artificial**
```
142 <220> FEATURE:
143 <223> OTHER INFORMATION: oligonucleotide
145 <400> SEQUENCE: 10
146 cttcagcacc gcggacagtg cctgtccatt cctgcacgac gtaccttcag caccgcggac   60
148 agtgcctgtc cattcctgca cgacgtacct tcagcaccgc ggacagtgcc tgtccattcc  120
150 tgcacgacgt ac                132
```

RAW SEQUENCE LISTING   DATE: 05/23/2001
PATENT APPLICATION: US/09/723,326   TIME: 10:32:47

Input Set : A:\09723326.txt
Output Set: C:\CRF3\05232001\I723326.raw

```
        153  <210> SEQ ID NO: 11
        154  <211> LENGTH: 116
        155  <212> TYPE: DNA
C-->    156  <213> ORGANISM: Artificial
        158  <220> FEATURE:
        159  <223> OTHER INFORMATION: oligonucleotide
        161  <400> SEQUENCE: 11
        162  cttcagcacc gcggacagtt gaggggactt tcccaggctt cagcaccgcg gacagttgag     60
        164  gggactttcc caggcttcag caccgcggac agttgagggg actttcccag gcgtac        116
        167  <210> SEQ ID NO: 12
        168  <211> LENGTH: 116
        169  <212> TYPE: DNA
C-->    170  <213> ORGANISM: Artificial
        172  <220> FEATURE:
        173  <223> OTHER INFORMATION: oligonucleotide
        175  <400> SEQUENCE: 12
        176  cttcagcacc gcttacagtt gaggggactt tcccaggctt cagcaccgct tacagttgag     60
        178  gggactttcc caggcttcag caccgcttac agttgagggg actttcccag gcgtac        116
        181  <210> SEQ ID NO: 13
        182  <211> LENGTH: 199
        183  <212> TYPE: DNA
C-->    184  <213> ORGANISM: Artificial
        186  <220> FEATURE:
        187  <223> OTHER INFORMATION: oligonucleotide
        189  <400> SEQUENCE: 13
        190  cttcagcacc gcggacagtt gacacgatca cctcccatta aggagagaga tctccttcag     60
        192  caccgcggac agttgacacg atcacctccc attaaggaga gagatctcct tcagcaccgc    120
        194  ggacagttga cacgatcacc tcccattaag gagagagatc tccttcagca ccgcggacag    180
        196  ttgaggggac tttcccagg                                                 199
        199  <210> SEQ ID NO: 14
        200  <211> LENGTH: 162
        201  <212> TYPE: DNA
C-->    202  <213> ORGANISM: Artificial
        204  <220> FEATURE:
        205  <223> OTHER INFORMATION: oligonucleotide
        207  <400> SEQUENCE: 14
        208  cttcagcacc gcggacagtt gagcttcggg gcttttgcac tcgtcccggc tctacttcag     60
        210  caccgcggac agttgagctt cggggctttt gcactcgtcc cggctctact tcagcaccgc    120
        212  ggacagttga gcttcggggc ttttgcactc gtcccggctc ta                       162
        215  <210> SEQ ID NO: 15
        216  <211> LENGTH: 52
        217  <212> TYPE: DNA
C-->    218  <213> ORGANISM: Artificial
        220  <220> FEATURE:
        221  <223> OTHER INFORMATION: oligonucleotide
        223  <400> SEQUENCE: 15
        224  ctggccttat ctccggctgc acgttgcctg ttggtgacta ataacacaat aa              52
        227  <210> SEQ ID NO: 16
        228  <211> LENGTH: 162
```

RAW SEQUENCE LISTING   DATE: 05/23/2001
PATENT APPLICATION: US/09/723,326   TIME: 10:32:47

Input Set : A:\09723326.txt
Output Set: C:\CRF3\05232001\I723326.raw

```
       229 <212> TYPE: DNA
C-->   230 <213> ORGANISM: Artificial
       232 <220> FEATURE:
       233 <223> OTHER INFORMATION: oligonucleotide
       235 <400> SEQUENCE: 16
       236 cttcagcacc gcggacagtt gacacgatca cctcccatta aggagagaga tctccttcag    60
       238 caccgcggac agttgacacg atcacctccc attaaggaga gagatctcct tcagcaccgc   120
       240 ggacagttga cacgatcacc tcccattaag gagagagatc tc                      162
```

VERIFICATION SUMMARY   DATE: 05/23/2001
PATENT APPLICATION: **US/09/723,326**   TIME: 10:32:48

Input Set : **A:\09723326.txt**
Output Set: **C:\CRF3\05232001\I723326.raw**

L:22 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:34 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:46 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:58 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:70 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:82 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:96 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:112 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:124 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:140 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:156 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:170 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:184 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:202 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:218 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:230 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16